

Figure 1 A

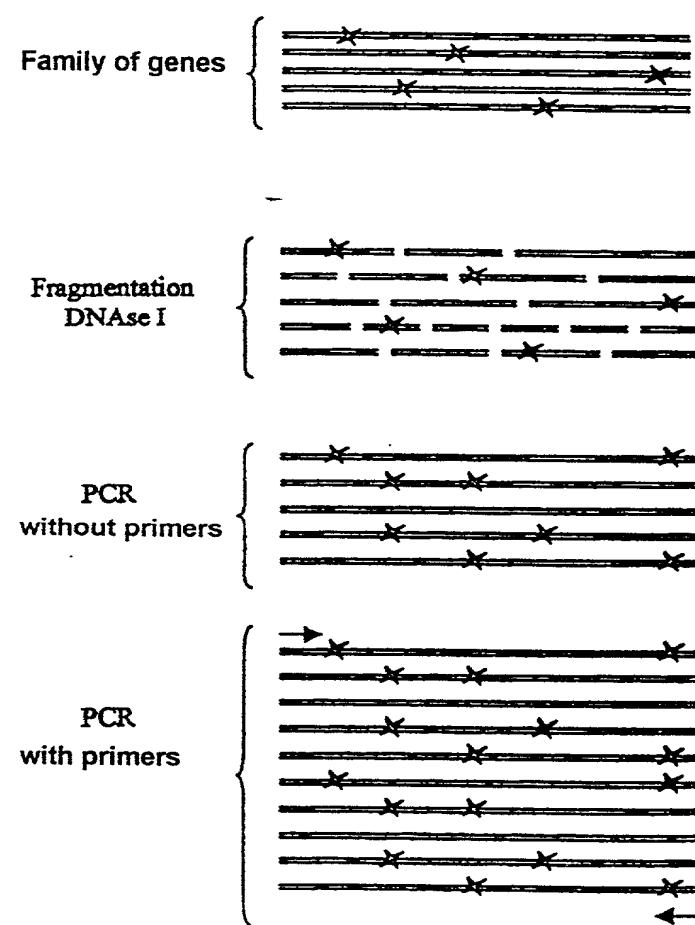


Figure 1 B

(Double-stranded process carried out, but illustrated here with a single strand)

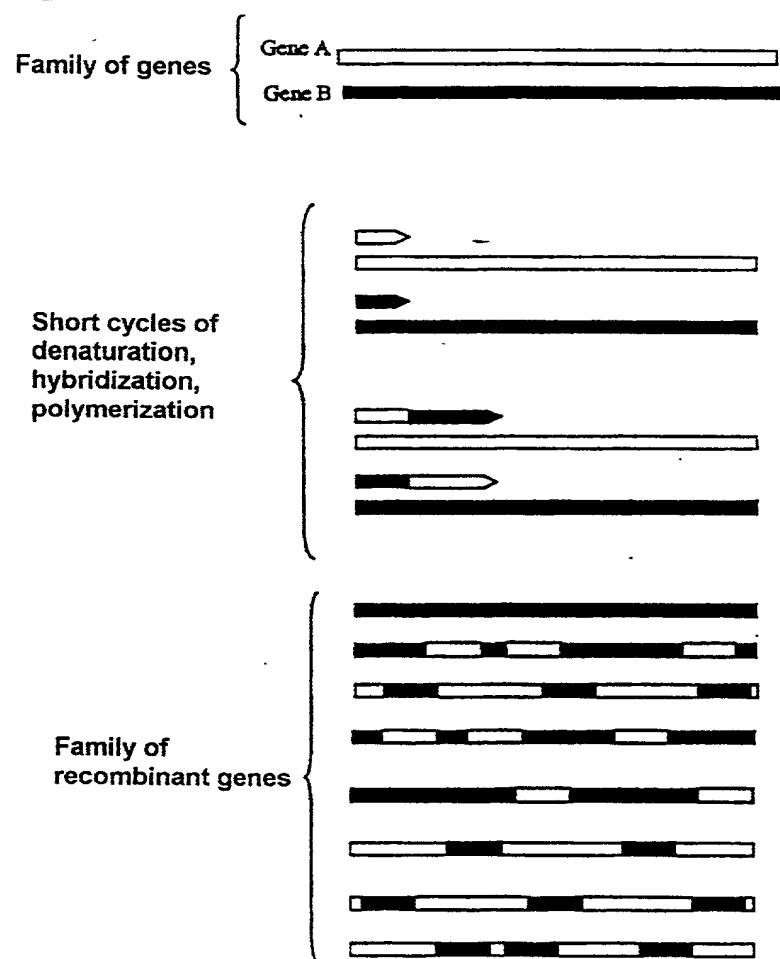


Fig. 2

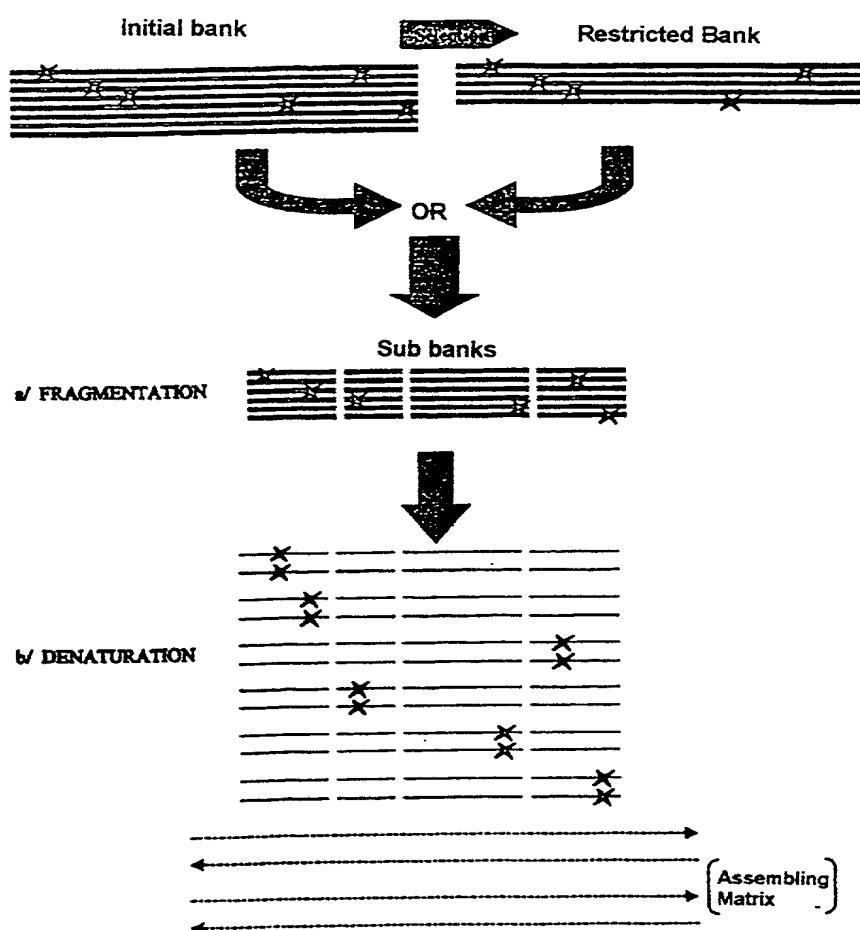


Fig. 2- continued

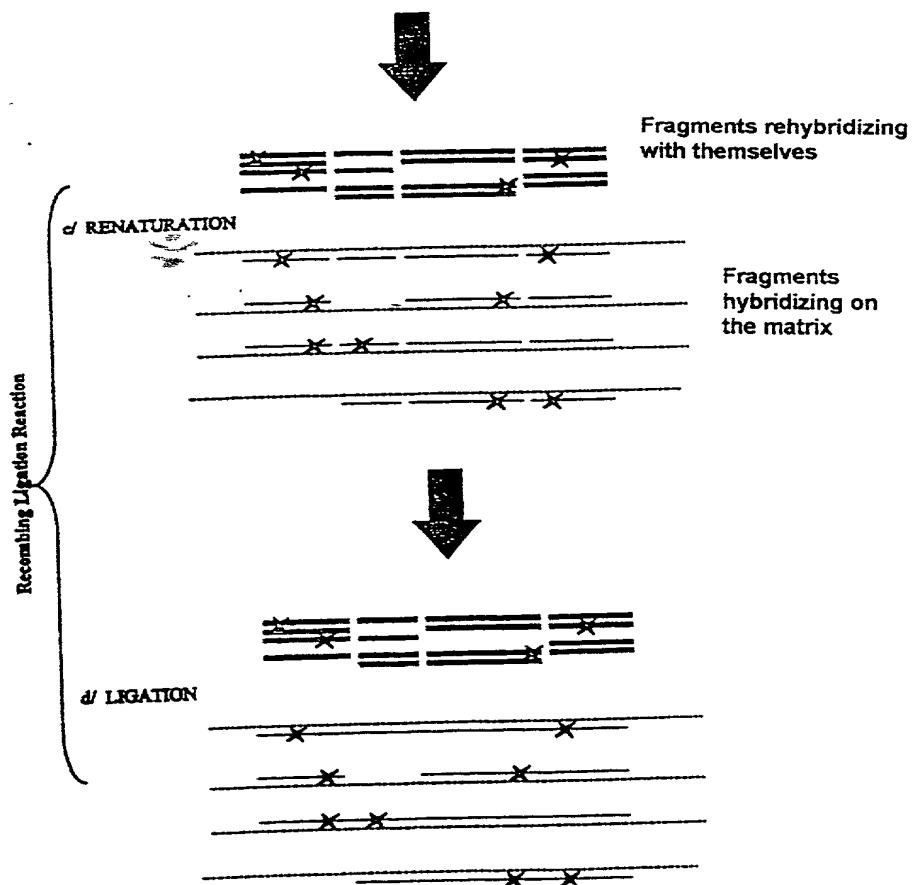
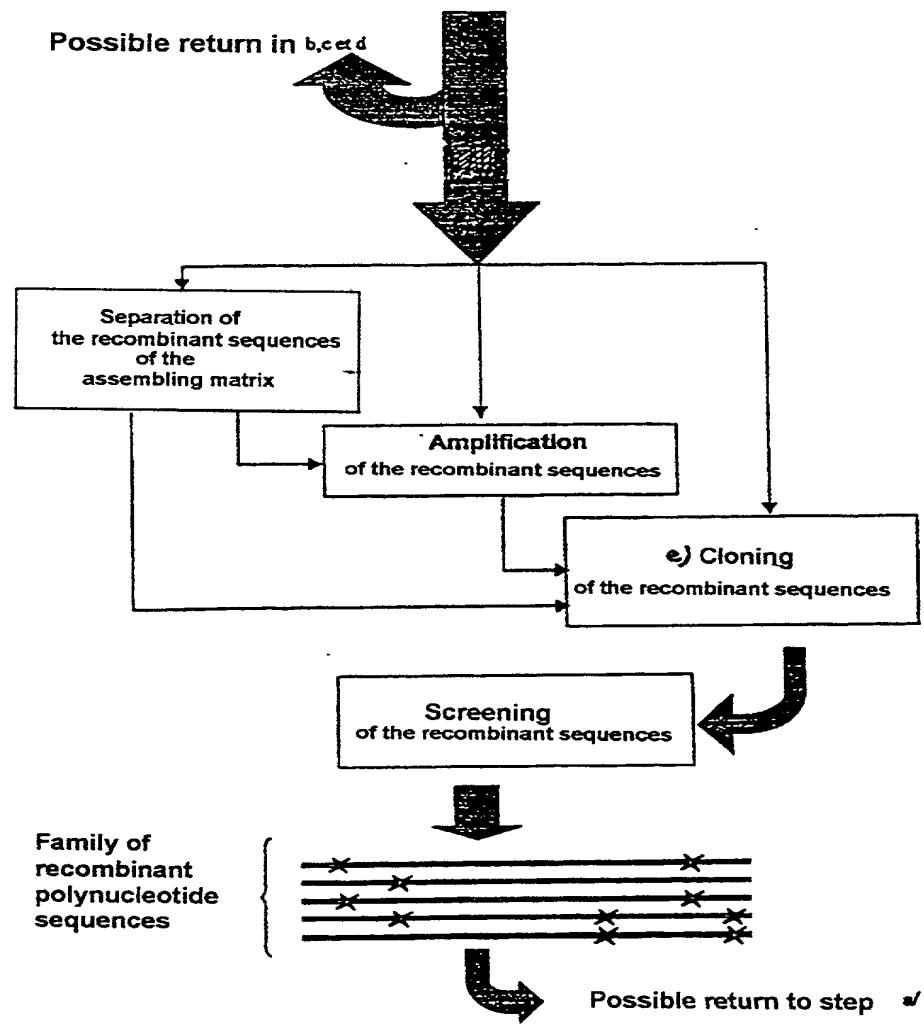


Fig. 2 continued



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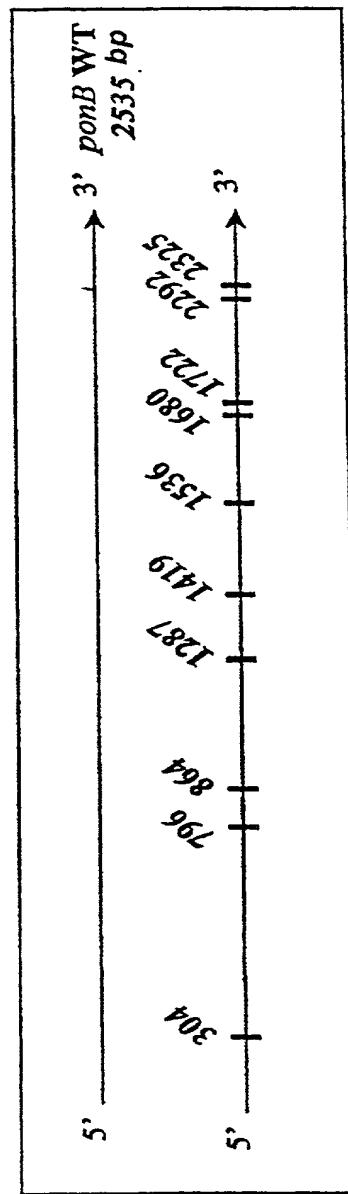


Figure 3: Position of the ten mutation zones (sites *Pvu* *II* and *Pst* *I*)

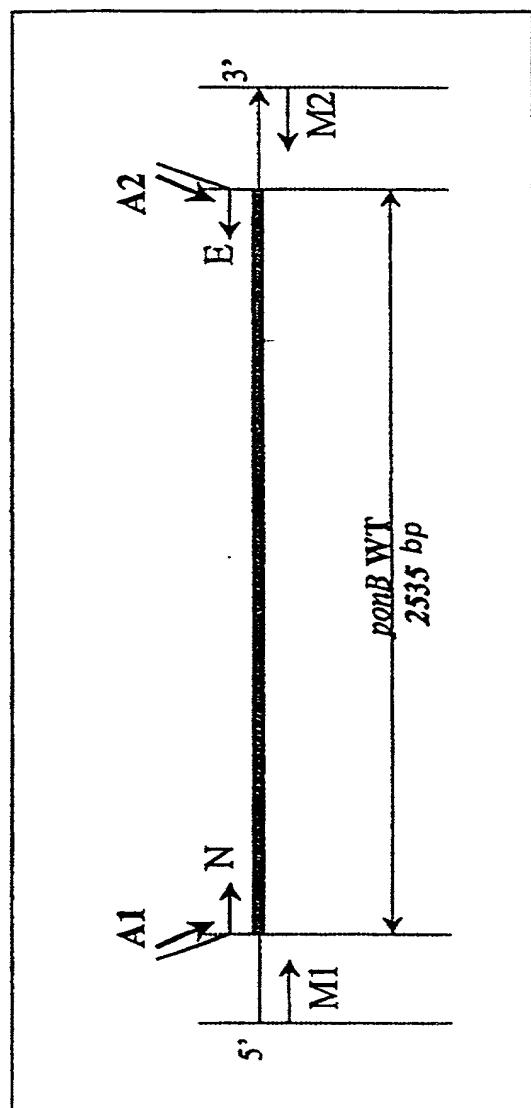


Figure 4 : Position of the primers used as compared to the sequence of the *pconB* gene

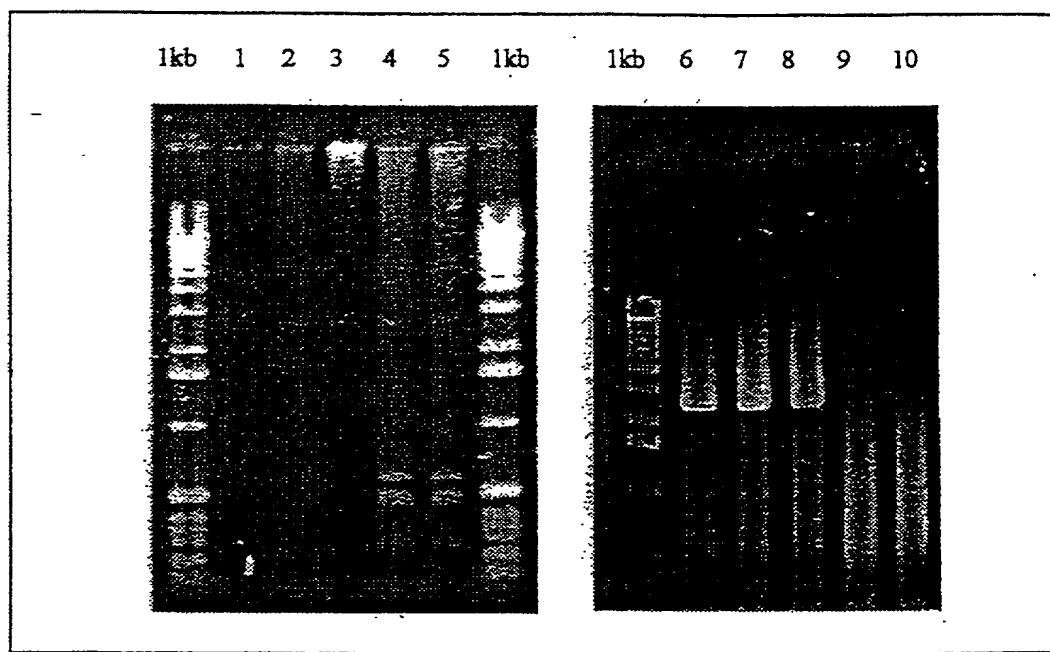


Fig. 5 : Migration of the RLR reactions and of the PCR amplifications of these reactions

<i>Tracks:</i>	<i>1/RLR 1</i>	<i>6/PCR RLR 1</i>
	<i>2/RLR 2</i>	<i>7/PCR RLR 2</i>
	<i>3/RLR 3</i>	<i>8/PCR RLR 3</i>
	<i>4/RLR 4</i>	<i>9/PCR RLR 4</i>
	<i>5/ RLR Control</i>	<i>10/PCR RLR Control</i>

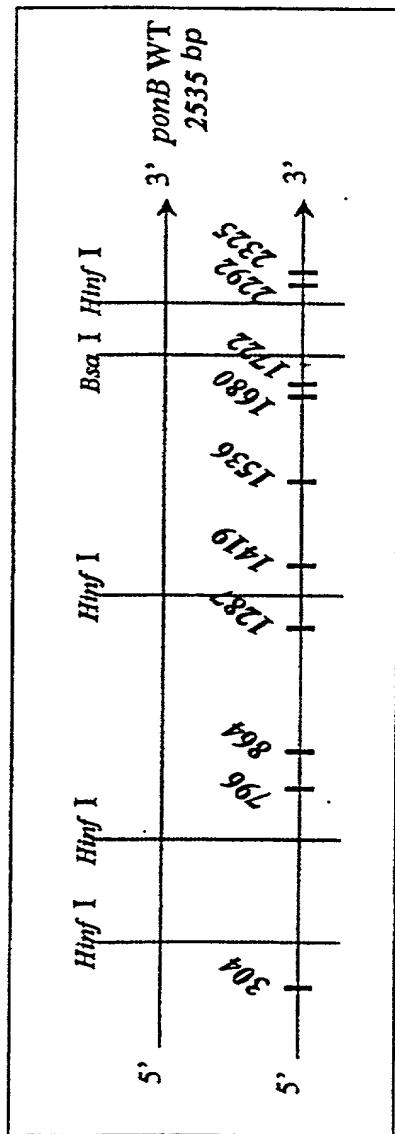


Figure 6: Position of the mutations as compared to the restriction fragments